OIPE #6 WO 215502

CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date: 8/29/2001

Serial Number: 09/763,957 ENTERED

Edited by:

Verified by: AV (STIC sta

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected: _____
- ☐ Other: _____

"Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING DATE: 08/29/2001
PATENT APPLICATION: US/09/763,957 TIME: 13:06:54

Input Set : A:\Pto.amc
Output Set: N:\CRF3\08292001\I763957.raw

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      5 <120> TITLE OF INVENTION: A NOVEL PLANT PROMOTER AND USES THEREFOR
      7 <130> FILE REFERENCE: 2209376/EJH
C-->  9 <140> CURRENT APPLICATION NUMBER: US/09/763,957
C--> 10 <141> CURRENT FILING DATE: 2001-02-28
     12 <150> PRIOR APPLICATION NUMBER: PP5572
     13 <151> PRIOR FILING DATE: 1998-08-31
     15 <160> NUMBER OF SEQ ID NOS: 9
     17 <170> SOFTWARE: PatentIn Ver. 2.0
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     39 cca tac ttt gat gga tgg aag gct tat gat caa aac ccc ttt cat ccc   210
     40 Pro Tyr Phe Asp Gly Trp Lys Ala Tyr Asp Gln Asn Pro Phe His Pro
     41                  30                  35                  40
     43 aca gat aat cct aac ggt gtt atg caa atg ggt ctt gct gag aat cag   258
     44 Thr Asp Asn Pro Asn Gly Val Met Gln Met Gly Leu Ala Glu Asn Gln
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     47 ctt acc tct gat ttg gtt gaa gat tgg ata ctg aac aac cct gaa gcc   306
     48 Leu Thr Ser Asp Leu Val Glu Asp Trp Ile Leu Asn Asn Pro Glu Ala
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     51 tcc att tgc act cca gaa gga ata aat gat ttc agg gcc ata gct aac   354
     52 Ser Ile Cys Thr Pro Glu Gly Ile Asn Asp Phe Arg Ala Ile Ala Asn
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     56 Phe Gln Asp Tyr His Gly Leu Ala Glu Phe Arg Asn Ala Val Ala Lys
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     64 Ile Val Met Ser Gly Gly Ala Thr Gly Ala His Glu Val Thr Ala Phe
     65             125                 130                 135
     67 tgt ttg gca gat ccc ggc gag gca ttc tta gtg ccc att ccc tat tat   546
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79 ttg gaa gat gcc tat gag aaa gca aga gag gat aac atc aga gta aag   690
80 Leu Glu Asp Ala Tyr Glu Lys Ala Arg Glu Asp Asn Ile Arg Val Lys
81                  190                 195                 200
83 ggt tta ctg atc acc aat cca tca aat cca tta ggc aca atc atg gac   738
84 Gly Leu Leu Ile Thr Asn Pro Ser Asn Pro Leu Gly Thr Ile Met Asp
85            205                 210                 215
87 aga aag aca ctg aga acc gtg gtg agc ttc atc aat gag aag cgt atc   786
88 Arg Lys Thr Leu Arg Thr Val Val Ser Phe Ile Asn Glu Lys Arg Ile
89          220                 225                 230
91 cac ctt gta tgt gat gaa ata tat gct gca aca gtt ttc agc caa ccc   834
92 His Leu Val Cys Asp Glu Ile Tyr Ala Ala Thr Val Phe Ser Gln Pro
93      235                 240                 245
95 ggt ttc ata agc ata gct gag ata tta gag gat gaa aca gac ata gag   882
96 Gly Phe Ile Ser Ile Ala Glu Ile Leu Glu Asp Glu Thr Asp Ile Glu
97 250                 255                 260                 265
99 tgt gac cgc aac ctc gta cac att gtt tat agt ctt tca aag gac atg   930
100 Cys Asp Arg Asn Leu Val His Ile Val Tyr Ser Leu Ser Lys Asp Met
101                 270                 275                 280
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116 Glu Arg Phe Leu Ala Glu Ser Ala Lys Arg Leu Ala Gln Arg Phe Arg
117 330                 335                 340                 345
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120 Val Phe Thr Gly Gly Leu Ala Lys Val Gly Ile Lys Cys Leu Gln Ser
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123 aat gct ggt cta ttt gtg tgg atg gat tta agg caa ctt ctc aaa aag   1218
124 Asn Ala Gly Leu Phe Val Trp Met Asp Leu Arg Gln Leu Leu Lys Lys
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139 att gct ttg caa cga atc cgc aac ttt gtg ctt caa aac aag gag gtc   1410
140 Ile Ala Leu Gln Arg Ile Arg Asn Phe Val Leu Gln Asn Lys Glu Val
141                 430                 435                 440
143 gtg gtg tct aat aag aaa cat tgt tgg cac agt aac ttg agg ctg agc   1458
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145             445                 450                 455
147 ctc aaa acc aga agg ttt gat gat atc acc atg tca cct cac tct ccc   1506
148 Leu Lys Thr Arg Arg Phe Asp Asp Ile Thr Met Ser Pro His Ser Pro
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151 cta cct cag tca cct atg gtt aaa gcc aca aat tgagtttgca tattcctctg 1559
152 Leu Pro Gln Ser Pro Met Val Lys Ala Thr Asn
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183 Ala Tyr Asp Gln Asn Pro Phe His Pro Thr Asp Asn Pro Asn Gly Val
185         35                  40                  45
187 Met Gln Met Gly Leu Ala Glu Asn Gln Leu Thr Ser Asp Leu Val Glu
189     50                  55                  60
191 Asp Trp Ile Leu Asn Asn Pro Glu Ala Ser Ile Cys Thr Pro Glu Gly
193 65                  70                  75                  80
195 Ile Asn Asp Phe Arg Ala Ile Ala Asn Phe Gln Asp Tyr His Gly Leu
197                 85                  90                  95
199 Ala Glu Phe Arg Asn Ala Val Ala Lys Phe Met Ala Arg Thr Arg Gly
201             100                 105                 110
203 Asn Arg Ile Thr Phe Asp Pro Asp Arg Ile Val Met Ser Gly Gly Ala
205         115                 120                 125
207 Thr Gly Ala His Glu Val Thr Ala Phe Cys Leu Ala Asp Pro Gly Glu
209     130                 135                 140
211 Ala Phe Leu Val Pro Ile Pro Tyr Tyr Pro Gly Phe Asp Arg Asp Leu
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217                 165                 170                 175
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229     210                 215                 220
231 Val Ser Phe Ile Asn Glu Lys Arg Ile His Leu Val Cys Asp Glu Ile
233 225                 230                 235                 240
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237                 245                 250                 255
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241             260                 265                 270
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265         355                 360                 365
267 Met Asp Leu Arg Gln Leu Leu Lys Lys Pro Thr Phe Asp Ser Glu Thr
269     370                 375                 380
271 Glu Leu Trp Lys Val Ile Ile His Glu Val Lys Ile Asn Val Ser Pro
273 385                 390                 395                 400
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277                 405                 410                 415
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281             420                 425                 430
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285         435                 440                 445
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310 atgcatttta tctagaagct gggaactgaa ccaaaaaaat agccagttga acaactgcag 240
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L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date